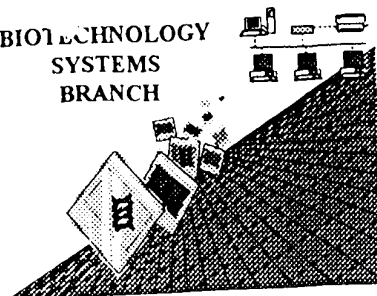


# **RAW SEQUENCE LISTING** **ERROR REPORT**

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/908,943  
Source: OIPÉ  
Date Processed by STIC: 7/30/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

## **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST 25. Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

# Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: <u>09/908,943</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 <u>    </u> Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor <b>after</b> creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <u>    </u> Invalid Line Length	The rules require that a line not <b>exceed</b> 72 characters in length. This includes white spaces.	
3 <u>    </u> Misaligned Amino Numbering	The numbering under each 5 <sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use <b>space characters</b> , instead.	
4 <u>    </u> Non-ASCII	The submitted file was <b>not</b> saved in ASCII(DOS) text, as <b>required</b> by the Sequence Rules. Please <b>ensure your subsequent submission is saved in ASCII text</b> .	
5 <u>    </u> Variable Length	Sequence(s) <u>    </u> contain n's or Xaa's representing more than one residue. <b>Per Sequence Rules, each n or Xaa can only represent a single residue.</b> Please present the <b>maximum</b> number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <u>    </u> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) <u>    </u> . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. <b>This applies to the mandatory &lt;220&gt;-&lt;223&gt; sections for Artificial or Unknown sequences.</b>	
7 <u>    </u> Skipped Sequences (OLD RULES)	Sequence(s) <u>    </u> missing. If intentional, please insert the following lines for <b>each</b> skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped  Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to <b>include</b> the skipped sequences.	
8 <u>    </u> Skipped Sequences (NEW RULES)	Sequence(s) <u>    </u> missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <u>    </u> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is <b>MANDATORY</b> if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <u>  </u> Invalid <213> Response	Per 1.823 of Sequence Rules, the only <b>valid</b> <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is <b>required</b> when <213> response is Unknown or is Artificial Sequence	
11 <u>  </u> Use of <220>	Sequence(s) <u>    </u> missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is <b>MANDATORY</b> if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <u>    </u> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 <u>    </u> Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is <b>not</b> used to represent any value not specifically a nucleotide.	

OIPE

## RAW SEQUENCE LISTING

DATE: 07/30/2001

PATENT APPLICATION: US/09/908,943

TIME: 11:01:18

Input Set : A:\00281.txt

Output Set: N:\CRF3\07302001\I908943.raw

pp 6-7

4 <110> APPLICANT: Yan, Riqiang  
 5 Tomasselli, Alfredo G.  
 6 Gurney, Mark E.  
 7 Emmons, Thomas L.  
 8 Bienkowski, Mike J.  
 9 Heinrikson, Robert L.  
 11 <120> TITLE OF INVENTION: SUBSTRATES AND ASSAYS FOR BETA-SECRETASE ACTIVITY  
 13 <130> FILE REFERENCE: 29915/00281  
 C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/908,943  
 C--> 16 <141> CURRENT FILING DATE: 2001-07-19  
 18 <160> NUMBER OF SEQ ID NOS: 197  
 20 <170> SOFTWARE: PatentIn Ver. 2.0  
 22 <210> SEQ ID NO: 1  
 23 <211> LENGTH: 2070  
 24 <212> TYPE: DNA  
 25 <213> ORGANISM: Homo sapiens  
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 30 ctgcccgtgc cccgggagac cgacgaagag cccgaggagc ccggccggag gggcagcttt 180  
 31 gtggagatgg tggacaacct gaggggcaag tgggggcagg gctactacgt ggagatgacc 240  
 32 gtgggcagcc ccccgagac gctcaacatc ctggtggata caggcagcag taactttgca 300  
 33 gtgggtgtg cccccaccc cttcctgcat cgtactacc agaggcagct gtccagcaca 360  
 34 taccgggacc tccggaagg tgtgtatgtg ccctacaccc agggcaagt ggaaggagg 420  
 35 ctgggcaccg acctggtaag catcccccat ggcccacag tcaactgtgc tgccaacatt 480  
 36 gctgccatca ctgaatcaga caagtcttc atcaacggct ccaactggga aggcacctg 540  
 37 gggctggcct atgctgagat tgccaggcct gacgactccc tggagccttt ctttgactct 600  
 38 ctggtaaagc agaccacgt tcccaacctc ttctccctgc acctttgtgg tgcgtgcttc 660  
 39 cccctcaacc agtctgaagt gctggcctct gtcggaggga gcatgatcat tggaggatc 720  
 40 gaccactcgc tgtacacagg cagtctctgg tatacaccca tccggcgga gtggtattat 780  
 41 gaggtcatca ttgtgcgggt ggagatcaat ggacaggatc tgaaaatgga ctgcaaggag 840  
 42 tacaactatg acaagagcat tgtggacagt ggcaccacca accttcgttt gcccaagaaa 900  
 43 gtgtttgaag ctgcagtcac atccatcaag gcagcctcct ccacggagaa gttccctgat 960  
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 45 ttcccagtc tctcactcta cctaattgggt gaggttacca accagtcctt ccgcatcacc 1080  
 46 atccttccgc agcaatacct gggccagtg gaagatgtgg ccacgtccca agacgactgt 1140  
 47 tacaagtttg ccatctcaca gtcacccag ggcactgtta tgggagctgt tatcatggag 1200  
 48 ggtctctacg ttgtctttga tggggcccga aaacgaattg gctttgctgt cagecgttgc 1260  
 49 catgtgcacg atgagttcag gacggcagcg gtggaaggcc cttttgtcac cttggacatg 1320  
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 52 cgtgcctcc gctgctgag ccagcagcat gatgactttg ctgatgacat ctccctgctg 1500  
 53 aagtgaggag gcccatgggc agaagataga gattcccttg gaccacacct ccgtggttca 1560  
 54 ctttggtcac aagtaggaga cacagatggc acctgtggcc agagcacctc aggacctcc 1620  
 55 ccaccacca aatgcctctg ccttgatgga gaaggaaaag gctggcaagg tgggttccag 1680  
 56 ggactgtacc ttaggaaac agaaaagaga agaaagaagc actctgctgg cgggaatact 1740  
 57 cttggtcacc tcaaatataa gtcgggaaat tctgctgctt gaaacttcag cctgaacct 1800

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001

TIME: 11:01:18

Input Set : A:\00281.txt

Output Set: N:\CRF3\07302001\I908943.raw

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59 gtactggcat cacacgcagg ttaccttggc gtgtgtccct gtggtaccct ggcagagaag 1920
60 agaccaagct tgtttccctg ctggccaaag tcagtaggag aggatgcaca gtttgctatt 1980
61 tgcttttagag acagggactg tataaacaag cctaacattg gtgcaaagat tgctcttga 2040
62 attaaaaaaaa aaaaaaaaaa aaaaaaaaaa 2070
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65 <211> LENGTH: 501
66 <212> TYPE: PRT
67 <213> ORGANISM: Homo sapiens
69 <400> SEQUENCE: 2
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71 1 5 10 15
73 Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser
74 20 25 30
76 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
77 35 40 45
79 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
80 50 55 60
82 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
83 65 70 75 80
85 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
86 85 90 95
88 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
89 100 105 110
91 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
92 115 120 125
94 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
95 130 135 140
97 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
98 145 150 155 160
100 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
101 165 170 175
103 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro Asp Asp
104 180 185 190
106 Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His Val Pro
107 195 200 205
109 Asn Leu Phe Ser Leu His Leu Cys Gly Ala Gly Phe Pro Leu Asn Gln
110 210 215 220
112 Ser Glu Val Leu Ala Ser Val Gly Gly Ser Met Ile Ile Gly Gly Ile
113 225 230 235 240
115 Asp His Ser Leu Tyr Thr Gly Ser Leu Trp Tyr Thr Pro Ile Arg Arg
116 245 250 255
118 Glu Trp Tyr Tyr Glu Val Ile Ile Val Arg Val Glu Ile Asn Gly Gln
119 260 265 270
121 Asp Leu Lys Met Asp Cys Lys Glu Tyr Asn Tyr Asp Lys Ser Ile Val
122 275 280 285
124 Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro Lys Lys Val Phe Glu Ala
125 290 295 300
127 Ala Val Lys Ser Ile Lys Ala Ala Ser Ser Thr Glu Lys Phe Pro Asp

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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001  
TIME: 11:01:18

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Output Set: N:\CRF3\07302001\I908943.raw

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130 Gly Phe Trp Leu Gly Glu Gln Leu Val Cys Trp Gln Ala Gly Thr Thr
131          325          330          335
133 Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met Gly Glu Val
134          340          345          350
136 Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln Tyr Leu Arg
137          355          360          365
139 Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr Lys Phe Ala
140          370          375          380
142 Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val Ile Met Glu
143 385          390          395          400
145 Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile Gly Phe Ala
146          405          410          415
148 Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala Ala Val Glu
149          420          425          430
151 Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr Asn Ile Pro
152          435          440          445
154 Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val Met Ala Ala
155          450          455          460
157 Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys Gln Trp
158 465          470          475          480
160 Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe Ala Asp Asp
161          485          490          495
163 Ile Ser Leu Leu Lys
164          500
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169 <212> TYPE: DNA
170 <213> ORGANISM: Homo sapiens
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175 ctgcggetgc cccgggagac cgacgaagag cccgaggagc ccggccggag gggcagcttt 180
176 gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacgt ggagatgacc 240
177 gtgggcagcc ccccgagac gctcaacatc ctggtggata caggcagcag taactttgca 300
178 gtgggtgctg cccccaccc ctctctgcat cgctactacc agaggcagct gtccagcaca 360
179 taccgggacc tccggaaggg tgtgtatgtg ccctacaccc agggcaagtg ggaaggggag 420
180 ctgggcaccg acctggttaag catcccccat ggccccaacg tcaactgtgc tgccaacatt 480
181 gctgccatca ctgaatcaga caagtctctc atcaacggct ccaactggga aggcattctg 540
182 gggctggcct atgctgagat tgccaggctt tgtggtgctg gcttccccct caaccagtct 600
183 gaagtgtctg cctctgtcgg agggagcatg atcattggag gtatcgacca ctgctgttac 660
184 acaggcagtc tctggtatac acccatccgg cgggagtggt attatgaggt gatcattgtg 720
185 cgggtggaga tcaatggaca ggatctgaaa atggactgca aggagtacaa ctatgacaag 780
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188 gagcagctgg tgtgctggca agcaggcacc accccttgga acattttccc agtcatctca 960
189 ctctacctaa tgggtgaggt taccaaccag tccttcgcga tcaccatcct tccgcagcaa 1020
190 tacctgoggc cagtggaaga tgtggccacg tcccaagacg actgttataa gtttgccatc 1080
191 tcacagtcac ccacgggcac tgttatggga gctgttatca tggagggctt ctacgttgct 1140

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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001

TIME: 11:01:18

Input Set : A:\00281.txt

Output Set: N:\CRF3\07302001\I908943.raw

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194 aacattccac agacagatga gtcaaccctc atgaccatag cctatgtcat ggctgccatc 1320
195 tgcgcctctc tcatgctgcc actctgcctc atgggtgtgtc agtggcgctg cctccgctgc 1380
196 ctgcgccagc agcatgatga ctttgcctgt gacatctccc tgctgaagtg aggaggccca 1440
197 tgggcagaag atagagattc ccttggaacca cacctccgtg gttcactttg gtcacaagta 1500
198 ggagacacag atggcacctg tggccagagc acctcaggac cctccccacc caccaaatgc 1560
199 ctctgccttg atggagaagg aaaaggtctg caaggtgggt tccagggaact gtacctgtag 1620
200 gaaacagaaa agagaagaaa gaagcactct gctggcggga atactcttgg tcacctcaaa 1680
201 tttaagtcgg gaaattctgc tgcttgaaac ttcagccctg aacctttgtc caccattcct 1740
202 ttaaattctc caaccctaaag tattcttctt ttcttagttt cagaagtact ggcacacac 1800
203 gcaggttacc ttggcgtgtg tccctgtggt acctggcag agaagagacc aagcttgttt 1860
204 cctgtctggc caaagtcagt aggagaggat gcacagtttg ctatttgctt tagagacagg 1920
205 gactgtataa acaagcctaa cattggtgca aagattgctt cttgaaaaaa aaaaaaa 1977

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207 &lt;210&gt; SEQ ID NO: 4

208 &lt;211&gt; LENGTH: 476

209 &lt;212&gt; TYPE: PRT

210 &lt;213&gt; ORGANISM: Homo sapiens

212 &lt;400&gt; SEQUENCE: 4

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216 Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser
217           20           25           30
219 Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp
220           35           40           45
222 Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val
223           50           55           60
225 Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr
226   65           70           75           80
228 Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser
229           85           90           95
231 Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr
232           100          105          110
234 Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val
235           115          120          125
237 Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp
238           130          135          140
240 Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala Asn Ile
241   145          150          155          160
243 Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser Asn Trp
244           165          170          175
246 Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Leu Cys Gly
247           180          185          190
249 Ala Gly Phe Pro Leu Asn Gln Ser Glu Val Leu Ala Ser Val Gly Gly
250           195          200          205
252 Ser Met Ile Ile Gly Gly Ile Asp His Ser Leu Tyr Thr Gly Ser Leu
253           210          215          220
255 Trp Tyr Thr Pro Ile Arg Arg Glu Trp Tyr Tyr Glu Val Ile Ile Val
256   225          230          235          240

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## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001

TIME: 11:01:18

Input Set : A:\00281.txt

Output Set: N:\CRF3\07302001\I908943.raw

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258 Arg Val Glu Ile Asn Gly Gln Asp Leu Lys Met Asp Cys Lys Glu Tyr
259                245                250                255
261 Asn Tyr Asp Lys Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu
262                260                265                270
264 Pro Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser
265                275                280                285
267 Ser Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val
268                290                295                300
270 Cys Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser
271 305                310                315                320
273 Leu Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile
274                325                330                335
277 Leu Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln
278                340                345                350
280 Asp Asp Cys Tyr Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val
281                355                360                365
283 Met Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala
284                370                375                380
286 Arg Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu
287 385                390                395                400
289 Phe Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu
290                405                410                415
292 Asp Cys Gly Tyr Asn Ile Pro Gln Thr Asp Glu Ser Thr Leu Met Thr
293                420                425                430
295 Ile Ala Tyr Val Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu
296                435                440                445
298 Cys Leu Met Val Cys Gln Trp Arg Cys Leu Arg Cys Leu Arg Gln Gln
299                450                455                460
301 His Asp Asp Phe Ala Asp Asp Ile Ser Leu Leu Lys
302 465                470                475
305 <210> SEQ ID NO: 5
306 <211> LENGTH: 14
307 <212> TYPE: PRT
308 <213> ORGANISM: Artificial Sequence
310 <220> FEATURE:
311 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic
312     peptide sequence
314 <400> SEQUENCE: 5
315 Lys Val Glu Ala Asn Tyr Glu Val Glu Gly Glu Arg Lys Lys
316 1                5                10
319 <210> SEQ ID NO: 6
320 <211> LENGTH: 15
321 <212> TYPE: PRT
322 <213> ORGANISM: Artificial Sequence
324 <220> FEATURE:
325 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic
326     peptide sequence
328 <400> SEQUENCE: 6
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<210> SEQ ID NO 190

<211> LENGTH: 13

<212> TYPE: PRT

<213> ORGANISM: synthetic peptide sequence

<400> SEQUENCE: 190

Ser Glu Ile Ser Tyr Glu Val Glu Phe Arg Trp Lys Lys  
1 5 10

*unatd - see item 10 on Enw  
summary  
sheet*



09/908,943

<210> SEQ ID NO 194

<211> LENGTH: 6806

<212> TYPE: DNA

<213> ORGANISM: fusion protein comprising a maltose binding protein with 125 amino

acids from APP C-terminus.

*invalid sequence - also, this is a DNA*

*sequence,  
not a  
PRT*

*See item 10 on Eric Lunnay  
sheet*

*sequence*

*FBI*

The alphabet 'N' has been detected in the Sequence Listing.  
Please check the listing to ensure the correct alphabet  
is used in the sequence. The <210> to <213> fields of  
the sequence using n or Naa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001

TIME: 11:01:19

Input Set : A:\00281.txt

Output Set: N:\CRF3\07302001\I908943.raw

L:15 M:270 C: Current Application Number differs, Replaced Current Application Number  
 L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
 L:432 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13  
 L:470 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15  
 L:494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16  
 L:518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17  
 L:542 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18  
 L:589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
 L:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
 L:692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
 L:725 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28  
 L:728 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28  
 L:922 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41  
 L:1039 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49  
 L:1058 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50  
 L:1077 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51  
 L:1096 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52  
 L:1115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53  
 L:1134 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54  
 L:1153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55  
 L:1172 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56  
 L:1191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57  
 L:1210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58  
 L:1229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59  
 L:1248 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60  
 L:1267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61  
 L:1286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62  
 L:1305 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63  
 L:1324 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64  
 L:1343 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65  
 L:1362 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66  
 L:1381 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:67  
 L:1400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68  
 L:1420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69  
 L:1439 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70  
 L:1458 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71  
 L:1477 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72  
 L:1496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73  
 L:1515 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74  
 L:1534 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75  
 L:1553 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:76  
 L:1572 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77  
 L:1591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78  
 L:1610 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79  
 L:1629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80  
 L:1653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81  
 L:1677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82  
 L:1701 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/908,943

DATE: 07/30/2001

TIME: 11:01:19

Input Set : A:\00281.txt

Output Set: N:\CRF3\07302001\I908943.raw

L:1725 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84  
L:1749 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85  
L:1773 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86  
L:1792 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:87  
L:4215 M:259 W: Allowed number of lines exceeded, <213> ORGANISM: